

Figure 1

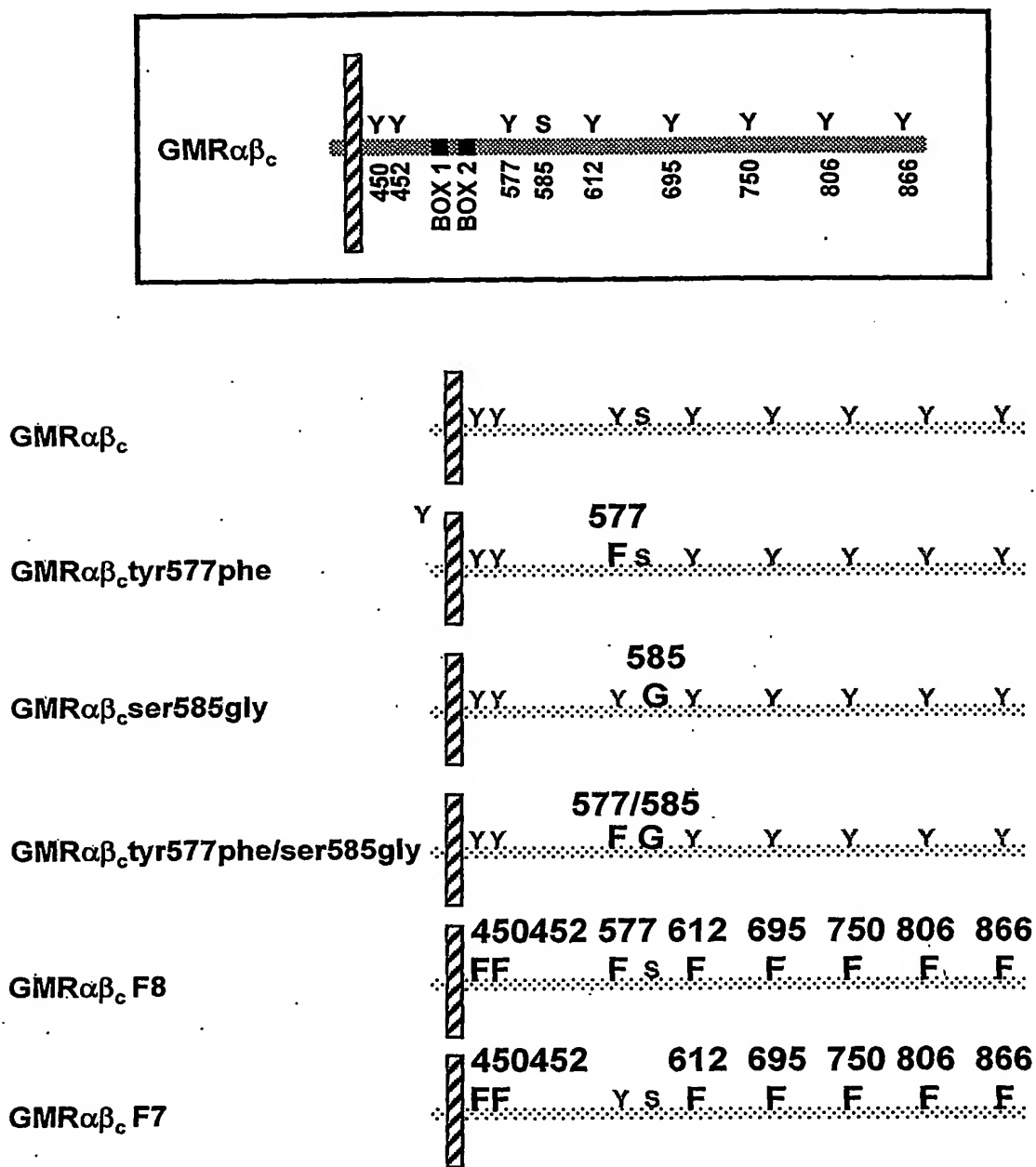


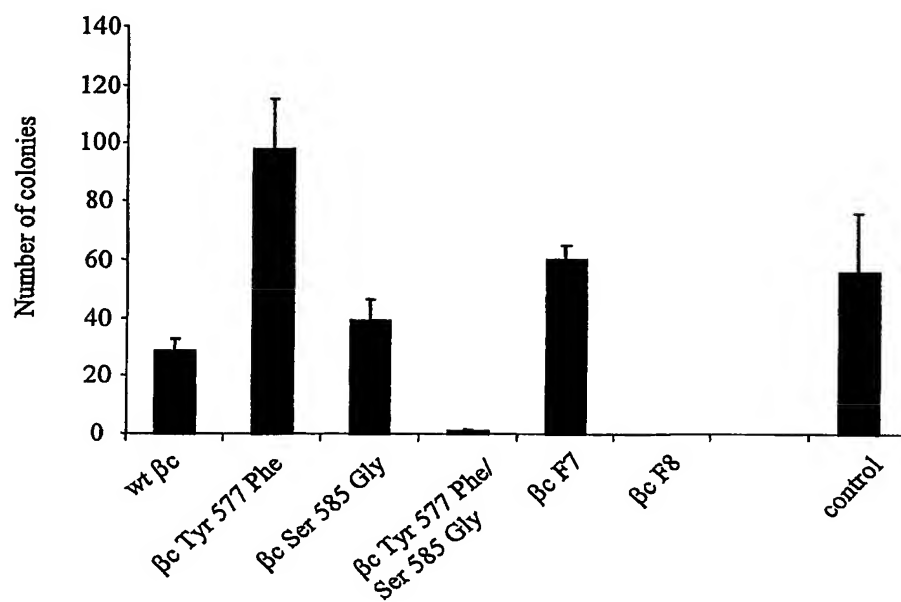
Figure 2

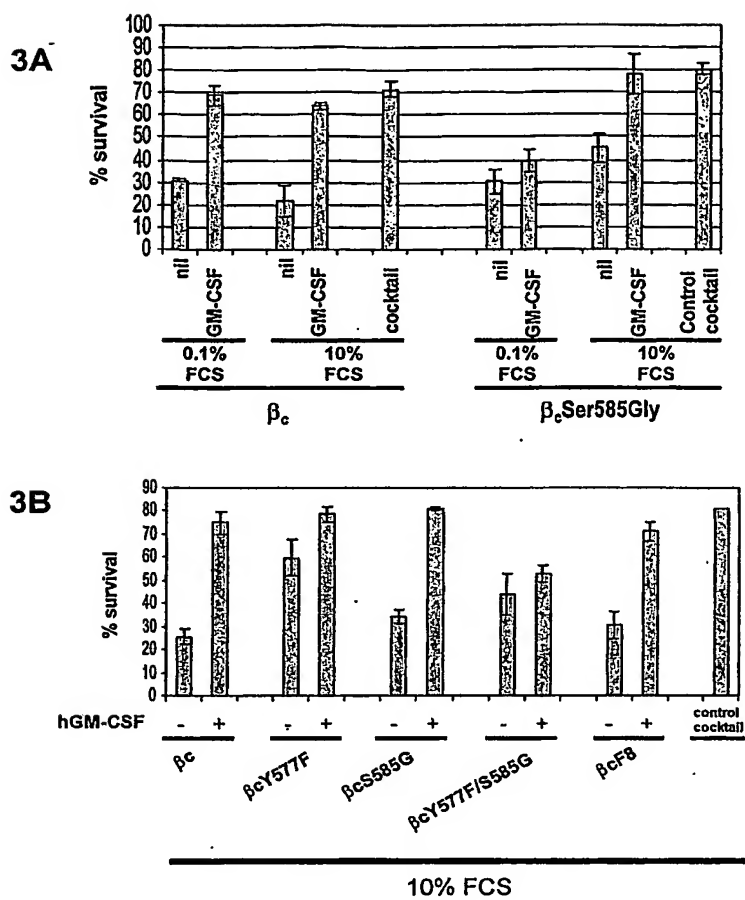
Figure 3

Figure 4

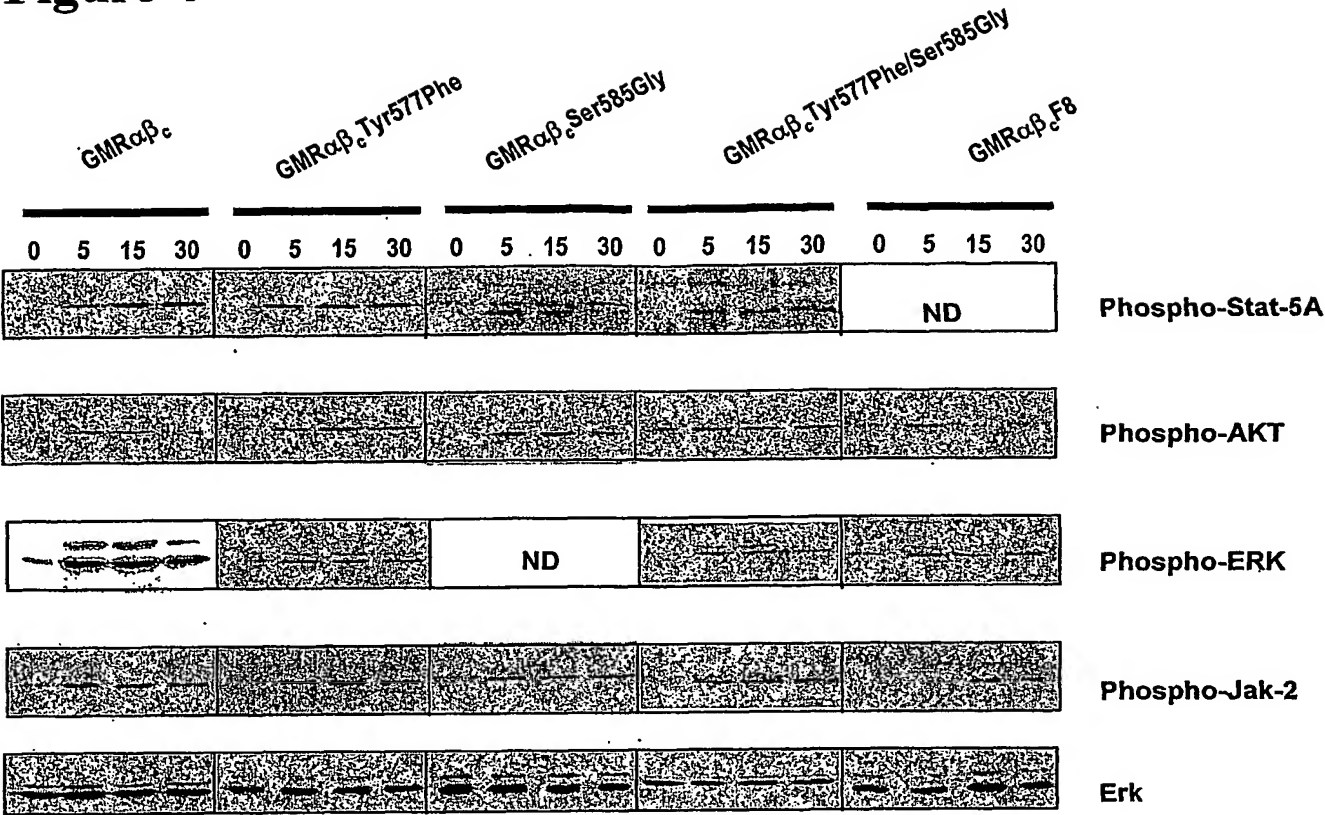


Figure 5

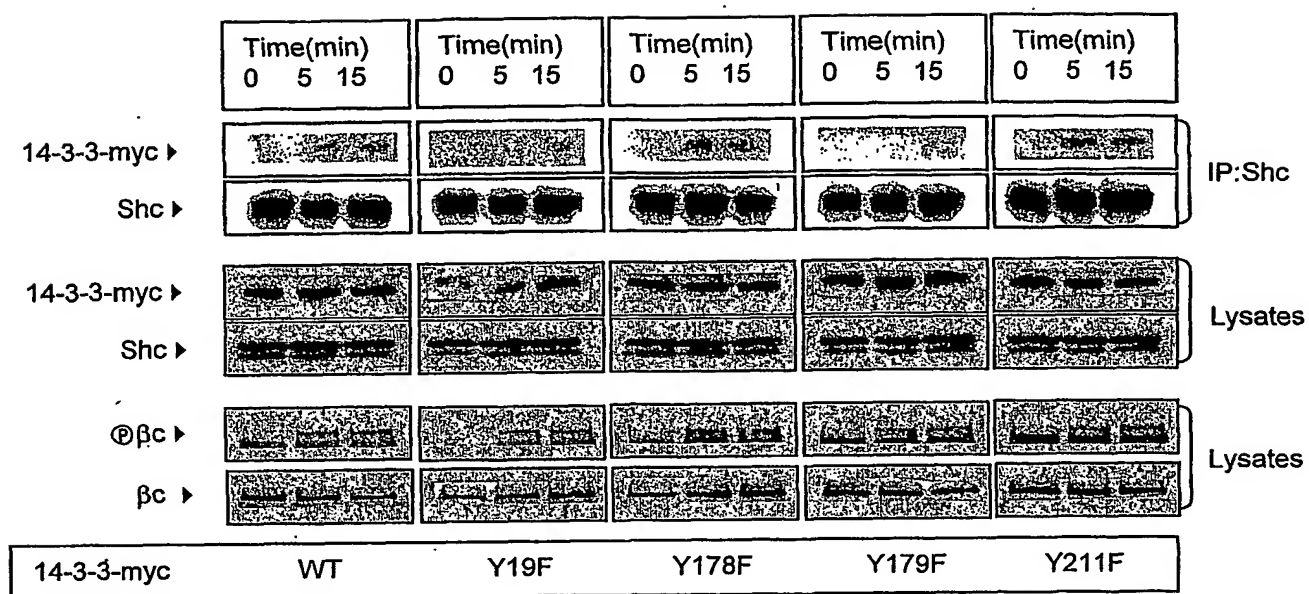
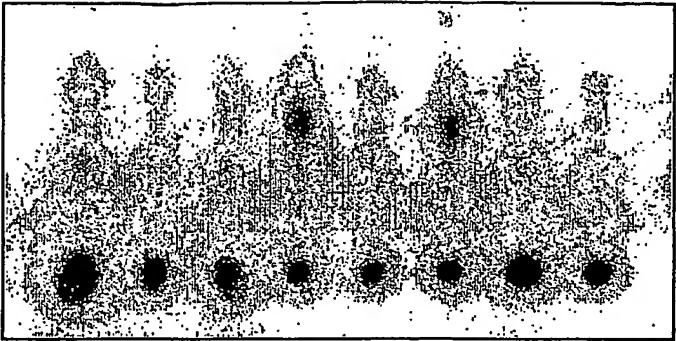


Figure 6

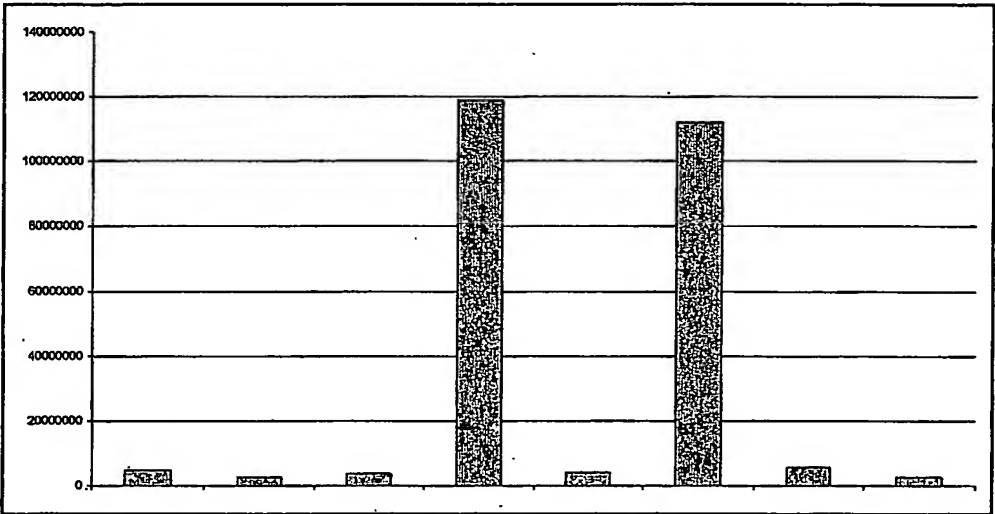
A

PIP →



GM-CSF	-	+	-	+	-	+	-	+
14-3-3-myc	-	-	WT	WT	Y178F	Y178F	Y179F	Y179F

B



GM-CSF	-	+	-	+	-	+	-	+
14-3-3-myc	-	-	WT	WT	Y178F	Y178F	Y179F	Y179F

Figure 7

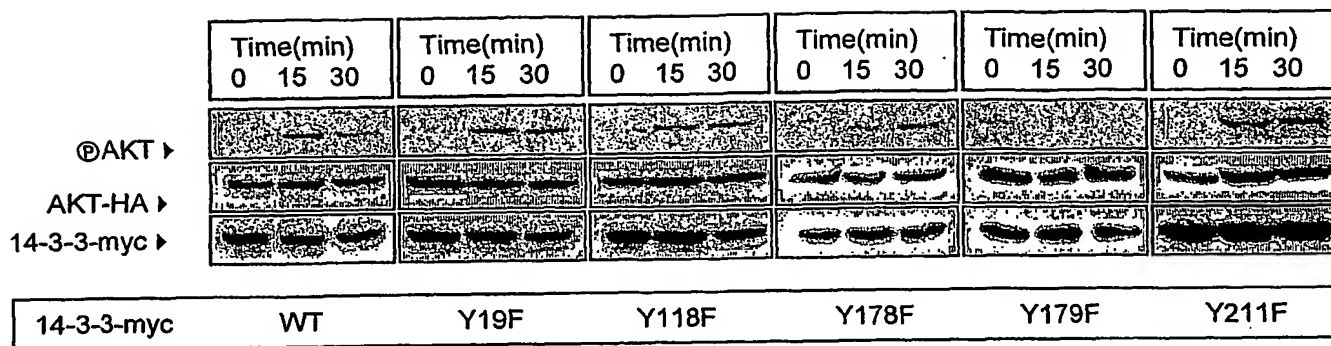
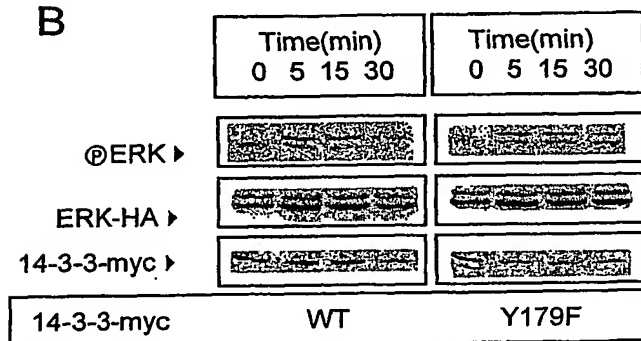
A**B**

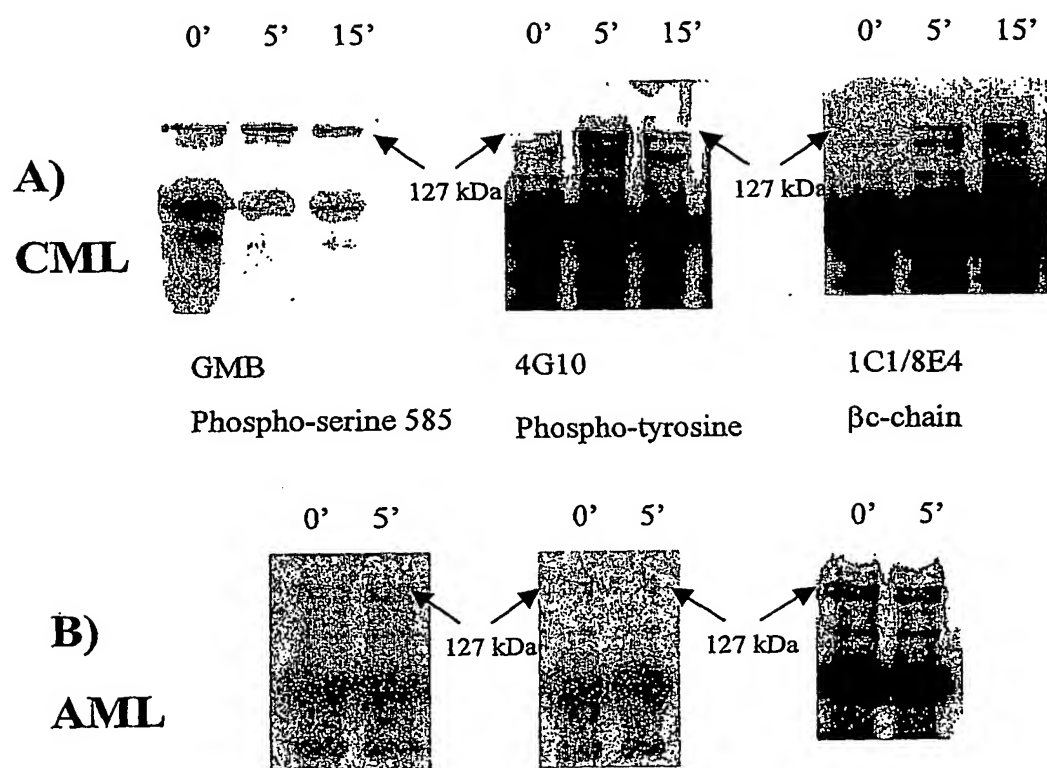
Figure 8

Figure 9

CYRB_HUMAN
 CYTOKINE RECEPTOR COMMON BETA CHAIN PRECURSOR (CDW131 ANTIGEN)
 Begin - 1, End - 897
 Seq: CYRB_HUMAN Length: 897 Fri Nov 17 13:50:29 2000 Check: 148

1	MVLAQGLLSM	ALLALCWERS	LAGAEETIPL	QTLRCYNDYT	SHITCRWADT
51	QDAQRLVNV	LIRRVNEDLL	EPVSCDLSDD	MPWSACPHPR	CVPRRCVIPC
101	QSFVVTDDY	FSFQPDRLG	TRLTVTLTQH	VQPPEPRDLQ	ISTDQDHFL
151	TWSVALGSPQ	SHWLSPGDLE	FEVVKRLQD	SWEDAAILLS	NTSQATLGPE
201	HLMPSSITYA	RVRTRLAPGS	RLSGRPSKWS	PEVCWDSQPG	DEAQPQNLEC
251	FFDGAAVLSC	SWEVKKEVAS	SVSFGLFYKP	SPDAGEEECS	PVLREGLGSL
301	HTRHHCQIPV	PDPATHGOYI	VSVQPRRAEK	HIKSSVNIQM	APPSLNVTKD
351	GDSYSLRWET	MKMRYEHIDH	TFEIQYRKDT	ATWKDSKTET	LQNAHSMALP
401	ALEPSTRYWA	RVRVRTSRTG	YNGIWSEWSE	ARSWDTESVL	PMWVLALIVI
451	FLTIAVLLAL	RFCGIYGYRL	RRKWEKIPN	PSKSHLFQNG	SAELWPPGSM
501	SAFTSGSPPH	QGPWGSRFPE	LEGVFPVGFG	DSEVSPLTIE	DPKHVCDPPS
551	GPDTTPAASD	LPTEQPPSPQ	PGPPAASHTP	EKQASSFDFN	GPYLGPPIHSR
601	SLPDILGQPE	PPQEGGSQKS	PPPGSLEYLC	LPAGGQVQLV	PLAQAMGPGQ
651	AVEVERRPSQ	GAAGSPSLES	GGGPAPPALG	PRVGGQDQKD	SPVAIPMSSG
701	DTEDPGVASG	YVSSADLVFT	PNSGASSVSL	VPSLGLPSDQ	TPSLCPGLAS
751	GPPGAPGPVK	SGFEGYVELP	PIEGRSPRSP	RNNPVPPEAK	SPVLNPGERP
801	ADVSPSTSPQ	EGLLVLQQVG	DYCFPLPGLP	GPLSLRSKPS	SPGPGPEIKN
851	LDQAFQVKKP	PGQAVPQVPV	IQLFKALKQQ	DYLSLPPWEV	NKPGEVC

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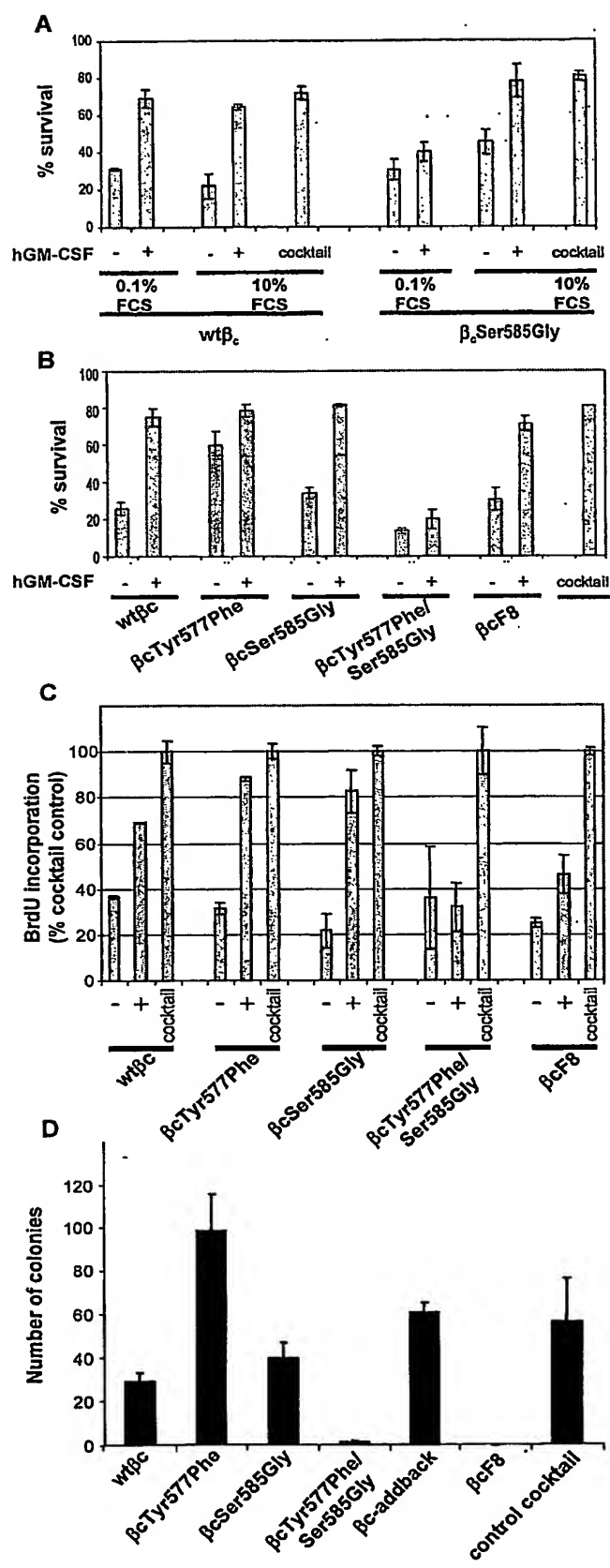
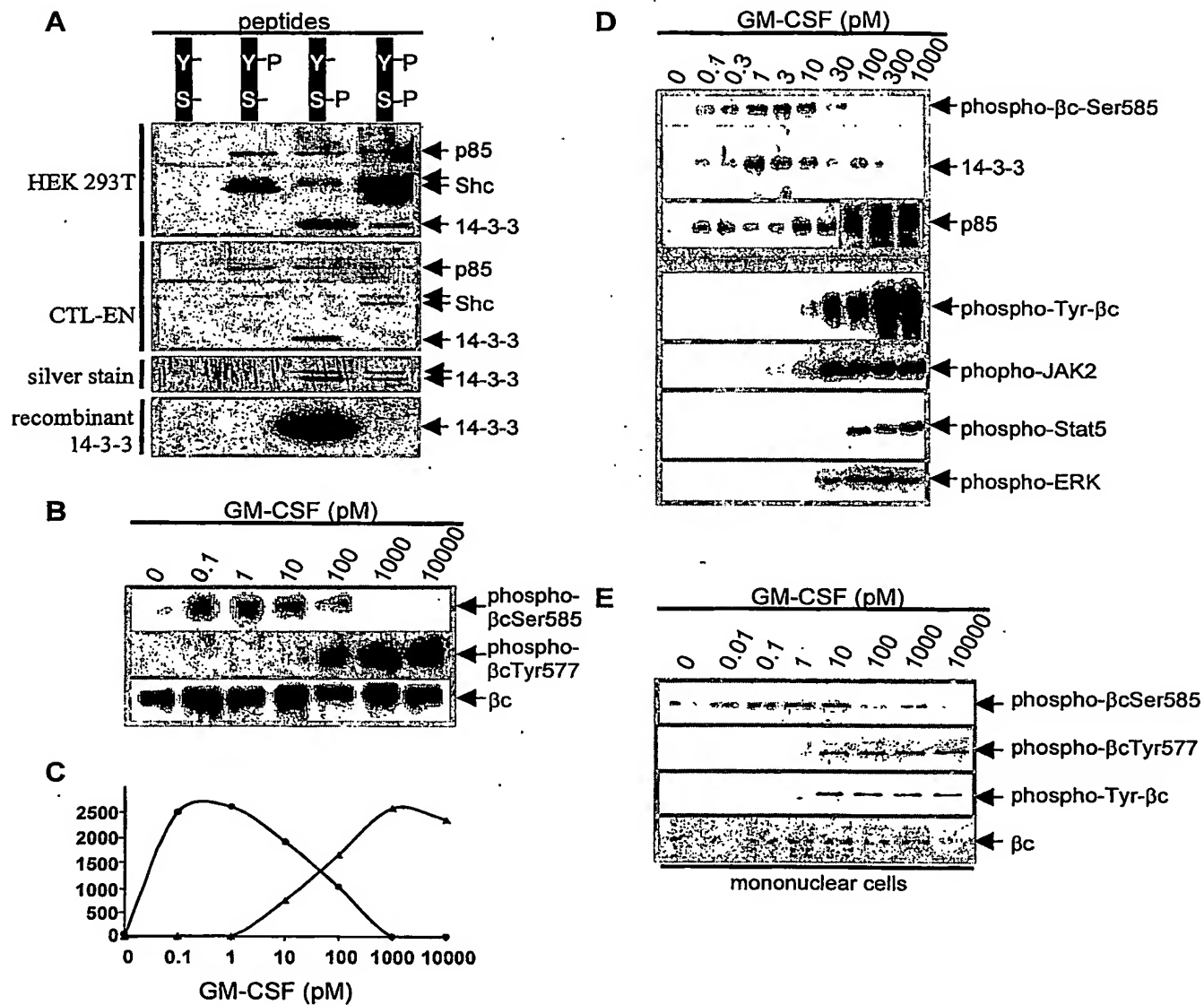
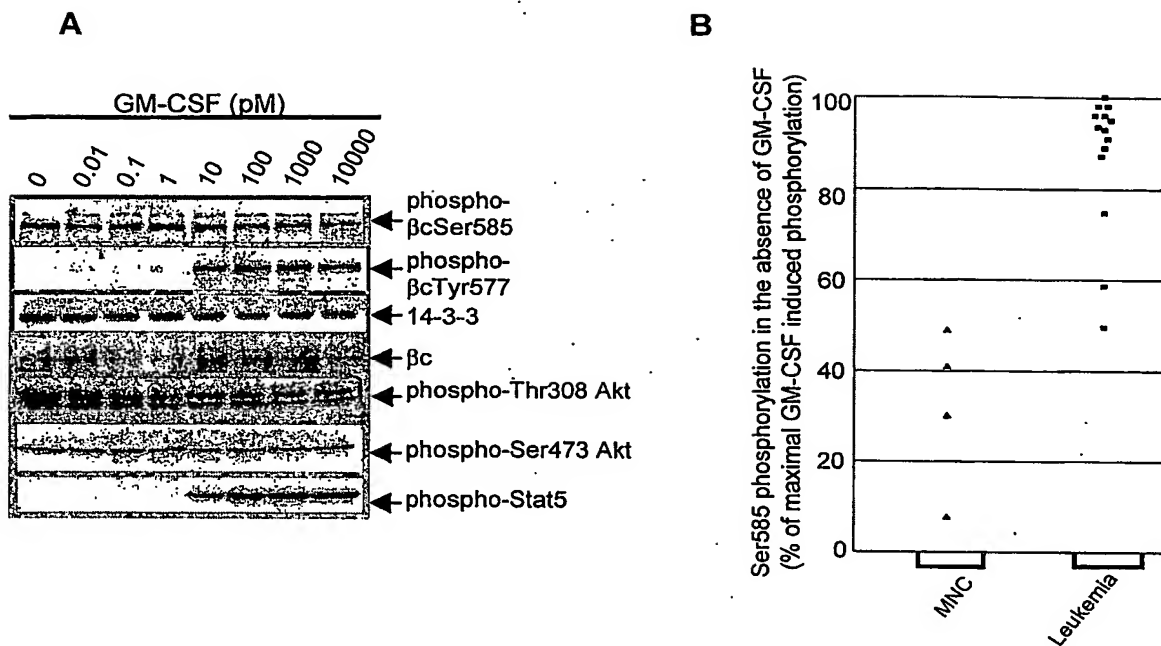
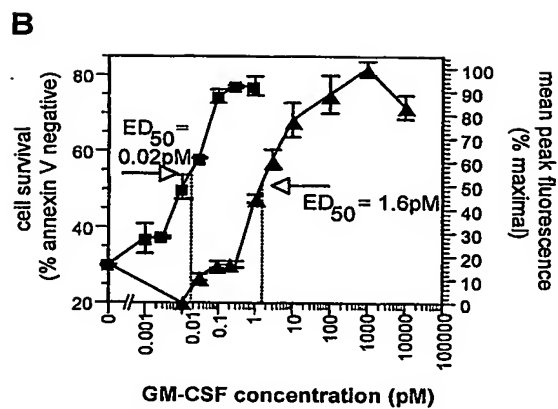
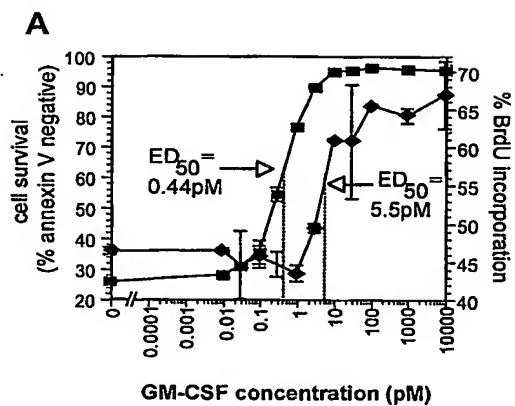
Figure 10

Figure 11

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Figure 12

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Figure 13

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Figure 15